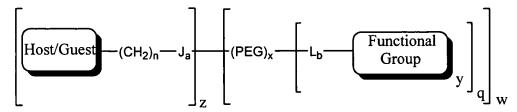
In the claims:

- 1-4. (Cancelled)
- 5. (Currently Amended) A composition comprising a cyclodextrin-containing polymer and a therapeutic agent and a complexing agent comprising at least one functional group and at least one host/guest moiety that forms an inclusion complex with a host/guest moiety of said polymer, wherein the polymer, the therapeutic agent, and the complexing agent are separate molecules.
- 6. (Previously Presented) A composition of claim 5, wherein said therapeutic agent is selected from an antibiotic, a steroid, a polynucleotide, small molecule pharmaceutical, a virus, a plasmid, a peptide, a peptide fragment, a chelating agent, a biologically active macromolecule, and mixtures thereof.
- 7. (Original) A composition of claim 6, wherein said therapeutic agent is a polynucleotide.

8-10. (Cancelled)

- 11. (Previously Presented) A composition of claim 5, wherein the host/guest of the complexing agent is selected from adamantyl, naphthyl, cholesterol, cyclodextrin, and mixtures thereof.
- 12. (Previously Presented) A composition of claim 5, wherein the complexing agent is a compound of the formula:



wherein

J is -NH-, $-C(=O)NH-CH_2)_{d^-}$, $-NH-C(=O)-(CH_2)_{d^-}$, $-CH_2SS-$, $-C(=O)O-(CH_2)_{e^-}O-P(=O)(O-CH_2)_{e^-}$

$$(CH_2)_e-Y)O-, \qquad , a peptide or polypeptide residue, or -NH-(C=O)-CH(R^1)-NH-; \\$$

Y is an additional host-guest functionality;

R¹ is -(CH₂)-CO₂H, an ester or salt thereof; or -(CH₂)_a-CONH₂;

PEG is $-O(CH_2CH_2O)_z$ -, where z varies from 2 to 500;

L is H, -NH, -NH-(C=O)-(CH₂)_e-(C=O)-CH₂-, -S(=O)₂-HC=CH-, -SS-, -C(=O)O-, or a carbohydrate residue;

a is 0 or 1;

b is 0 or 1;

d ranges from 0 to 6;

e ranges from 1 to 6;

n ranges from 0 to 6;

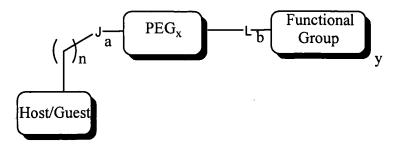
q ranges from 1 to 5;

w ranges from 1 to 5;

y is 1; and

x is 0 or 1.

13. (Previously Presented) A composition of claim 5, wherein the complexing agent is a compound of the formula:



wherein

J is -NH-, -C(=O)NH-CH₂)_d-, -NH-C(=O)-(CH₂)_d-, -CH₂SS-, -C(=O)O-(CH₂)_e-O-P(=O)(O-(CH₂)_e-O-P(=O)

, a peptide or polypeptide residue, or

-NH-(C=O)-CH(R¹)-NH-(C=O)-CH(R¹)-NH-;

Y is an additional host-guest functionality;

R¹ is -(CH₂)-CO₂H, an ester or salt thereof; or -(CH₂)_a-CONH₂;

PEG is $-O(CH_2CH_2O)_z$, where z varies from 2 to 500;

L is H, -NH, -NH-(C=O)-(CH₂)_e-(C=O)-CH₂-, -S(=O)₂-HC=CH-, -SS-, -C(=O)O-, or a carbohydrate residue;

a is 0 or 1;

b is 0 or 1;

d ranges from 0 to 6;

e ranges from 1 to 6;

n ranges from 0 to 6;

y is 1; and

x is 0 or 1.

(Previously Presented) A composition of claim 5, wherein the at least one functional 14. group includes a group selected from a ligand, a nuclear localization signal, an endosomal release peptide, an endosomal release polymer, or a membrane permeabilization agent.

- 15. (Previously Presented) A composition of claim 5, wherein the at least one functional group includes a moiety that increases the solubility of the composition under biological conditions relative to a composition of the polymer and therapeutic agent alone.
- 16. (Previously Presented) A composition of claim 5, wherein the at least one functional group includes a moiety that stabilizes the composition under biological conditions relative to a composition of the polymer and therapeutic agent alone.
- 17. (Previously Presented) A composition of claim 5, wherein the at least one functional group includes a therapeutic agent reversibly bound to the complexing agent.
- 18. (Previously Presented) A composition of claim 5, wherein the complexing agent further comprises a spacer group.
- 19. (Previously Presented) A composition of claim 5, wherein the polymer comprises a host moiety that forms an inclusion complex with a guest moiety of the complexing agent.
- 20. (Previously Presented) A composition of claim 5, wherein the polymer comprises a guest moiety that forms an inclusion complex with a host moiety of the complexing agent.
- 21. (Previously Presented) A composition of claim 20, wherein the guest moiety is an adamantyl group and the host moiety is a cyclodextrin moiety.